

SEQUENCE LISTING

<110> Matti Sallberg
Catharina Hultgren

<120> VACCINES CONTAINING RIBAVIRIN AND
METHODS OF USE THEREOF

<130> TRIPEP.23AUS2

<150> 09/705,547

<151> 2000-11-03

<150> 60/229,175

<151> 2000-08-29

<150> 60/225,767

<151> 2000-08-17

<160> 49

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 3011

<212> PRT

<213> Artificial Sequence

<220>

<223> Hepatitis C virus sequence

<400> 1

Met	Ser	Thr	Asn	Pro	Lys	Pro	Gln	Arg	Lys	Thr	Lys	Arg	Asn	Thr	Asn
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Arg	Arg	Pro	Gln	Asp	Val	Lys	Phe	Pro	Gly	Gly	Gly	Gln	Ile	Val	Gly
			20					25					30		
Gly	Val	Tyr	Leu	Leu	Pro	Arg	Arg	Gly	Pro	Arg	Leu	Gly	Val	Arg	Ala
		35				40					45				
Thr	Arg	Lys	Thr	Ser	Glu	Arg	Ser	Gln	Pro	Arg	Gly	Arg	Arg	Gln	Pro
	50					55				60					
Ile	Pro	Lys	Ala	Arg	Arg	Pro	Glu	Gly	Arg	Thr	Trp	Ala	Gln	Pro	Gly
65				70					75					80	
Tyr	Pro	Trp	Pro	Leu	Tyr	Gly	Asn	Glu	Gly	Cys	Gly	Trp	Ala	Gly	Trp
			85				90						95		
Leu	Leu	Ser	Pro	Arg	Gly	Ser	Arg	Pro	Ser	Trp	Gly	Pro	Thr	Asp	Pro
			100				105						110		
Arg	Arg	Arg	Ser	Arg	Asn	Leu	Gly	Lys	Val	Ile	Asp	Thr	Leu	Thr	Cys
		115				120					125				
Gly	Phe	Ala	Asp	Leu	Met	Gly	Tyr	Ile	Pro	Leu	Val	Gly	Ala	Pro	Leu
	130					135				140					
Gly	Gly	Ala	Ala	Arg	Ala	Leu	Ala	His	Gly	Val	Arg	Val	Leu	Glu	Asp
145				150					155					160	
Gly	Val	Asn	Tyr	Ala	Thr	Gly	Asn	Leu	Pro	Gly	Cys	Ser	Phe	Ser	Ile

Phe	Leu	Leu	Ala	Leu	Leu	Ser	Cys	Leu	Thr	Val	Pro	Ala	Ser	Ala	Tyr
Gln	Val	Arg	Asn	Ser	Ser	Gly	Leu	Tyr	His	Val	Thr	Asn	Asp	Cys	Pro
Asn	Ser	Ser	Val	Val	Tyr	Glu	Ala	Ala	Asp	Ala	Ile	Leu	His	Thr	Pro
Gly	Cys	Val	Pro	Cys	Val	Arg	Glu	Gly	Asn	Ala	Ser	Arg	Cys	Trp	Val
Ala	Val	Thr	Pro	Thr	Val	Ala	Thr	Arg	Asp	Gly	Lys	Leu	Pro	Thr	Thr
Gln	Leu	Arg	Arg	His	Ile	Asp	Leu	Leu	Val	Gly	Ser	Ala	Thr	Leu	Cys
Ser	Ala	Leu	Tyr	Val	Gly	Asp	Leu	Cys	Gly	Ser	Val	Phe	Leu	Val	Gly
Gln	Leu	Phe	Thr	Phe	Ser	Pro	Arg	His	His	Trp	Thr	Thr	Gln	Asp	Cys
Asn	Cys	Ser	Ile	Tyr	Pro	Gly	His	Ile	Thr	Gly	His	Arg	Met	Ala	Trp
Asn	Met	Met	Met	Asn	Trp	Ser	Pro	Thr	Ala	Ala	Leu	Val	Val	Ala	Gln
Leu	Leu	Arg	Ile	Pro	Gln	Ala	Ile	Met	Asp	Met	Ile	Ala	Gly	Ala	His
Trp	Gly	Val	Leu	Ala	Gly	Ile	Lys	Tyr	Phe	Ser	Met	Val	Gly	Asn	Trp
Ala	Lys	Val	Leu	Val	Val	Leu	Leu	Leu	Phe	Ala	Gly	Val	Asp	Ala	Glu
Thr	His	Val	Thr	Gly	Gly	Asn	Ala	Gly	Arg	Thr	Thr	Ala	Gly	Leu	Val
Gly	Leu	Leu	Thr	Pro	Gly	Ala	Lys	Gln	Asn	Ile	Gln	Leu	Ile	Asn	Thr
Asn	Gly	Ser	Trp	His	Ile	Asn	Ser	Thr	Ala	Leu	Asn	Cys	Asn	Glu	Ser
Leu	Asn	Thr	Gly	Trp	Leu	Ala	Gly	Leu	Phe	Tyr	Gln	His	Lys	Phe	Asn
Ser	Ser	Gly	Cys	Pro	Glu	Arg	Leu	Ala	Ser	Cys	Arg	Arg	Leu	Thr	Asp
Phe	Ala	Gln	Gly	Trp	Gly	Pro	Ile	Ser	Tyr	Ala	Asn	Gly	Ser	Gly	Leu
Asp	Glu	Arg	Pro	Tyr	Cys	Trp	His	Tyr	Pro	Pro	Arg	Pro	Cys	Gly	Ile
Val	Pro	Ala	Lys	Ser	Val	Cys	Gly	Pro	Val	Tyr	Cys	Phe	Thr	Pro	Ser
Pro	Val	Val	Val	Gly	Thr	Thr	Asp	Arg	Ser	Gly	Ala	Pro	Thr	Tyr	Ser
Trp	Gly	Ala	Asn	Asp	Thr	Asp	Val	Phe	Val	Leu	Asn	Asn	Thr	Arg	Pro
Pro	Leu	Gly	Asn	Trp	Phe	Gly	Cys	Thr	Trp	Met	Asn	Ser	Thr	Gly	Phe
Thr	Lys	Val	Cys	Gly	Ala	Pro	Pro	Cys	Val	Ile	Gly	Gly	Val	Gly	Asn
Asn	Thr	Leu	Leu	Cys	Pro	Thr	Asp	Cys	Phe	Arg	Lys	Tyr	Pro	Glu	Ala
Thr	Tyr	Ser	Arg	Cys	Gly	Ser	Gly	Pro	Arg	Ile	Thr	Pro	Arg	Cys	Met

Val	Asp	Tyr	Pro	Tyr	Arg	Leu	Trp	His	Tyr	Pro	Cys	Thr	Ile	Asn	Tyr	610	615	620
Thr	Ile	Phe	Lys	Val	Arg	Met	Tyr	Val	Gly	Gly	Val	Glu	His	Arg	Leu	625	630	635
Glu	Ala	Ala	Cys	Asn	Trp	Thr	Arg	Gly	Glu	Arg	Cys	Asp	Leu	Glu	Asp	645	650	655
Arg	Asp	Arg	Ser	Glu	Leu	Ser	Pro	Leu	Leu	Leu	Ser	Thr	Thr	Gln	Trp	660	665	670
Gln	Val	Leu	Pro	Cys	Ser	Phe	Thr	Thr	Leu	Pro	Ala	Leu	Ser	Thr	Gly	675	680	685
Leu	Ile	His	Leu	His	Gln	Asn	Ile	Val	Asp	Val	Gln	Tyr	Leu	Tyr	Gly	690	695	700
Val	Gly	Ser	Ser	Ile	Ala	Ser	Trp	Ala	Ile	Lys	Trp	Glu	Tyr	Val	Val	705	710	715
Leu	Leu	Phe	Leu	Leu	Leu	Ala	Asp	Ala	Arg	Val	Cys	Ser	Cys	Leu	Trp	725	730	735
Met	Met	Leu	Leu	Ile	Ser	Gln	Ala	Glu	Ala	Ala	Leu	Glu	Asn	Leu	Val	740	745	750
Ile	Leu	Asn	Ala	Ala	Ser	Leu	Ala	Gly	Thr	His	Gly	Leu	Val	Ser	Phe	755	760	765
Leu	Val	Phe	Phe	Cys	Phe	Ala	Trp	Tyr	Leu	Lys	Gly	Arg	Trp	Val	Pro	770	775	780
Gly	Ala	Val	Tyr	Ala	Leu	Tyr	Gly	Met	Trp	Pro	Leu	Leu	Leu	Leu	Leu	785	790	795
Leu	Ala	Leu	Pro	Gln	Arg	Ala	Tyr	Ala	Leu	Asp	Thr	Glu	Val	Ala	Ala	805	810	815
Ser	Cys	Gly	Gly	Val	Val	Leu	Val	Gly	Leu	Met	Ala	Leu	Thr	Leu	Ser	820	825	830
Pro	Tyr	Tyr	Lys	Arg	Tyr	Ile	Ser	Trp	Cys	Met	Trp	Trp	Leu	Gln	Tyr	835	840	845
Phe	Leu	Thr	Arg	Val	Glu	Ala	Gln	Leu	His	Val	Trp	Val	Pro	Pro	Leu	850	855	860
Asn	Val	Arg	Gly	Gly	Arg	Asp	Ala	Val	Ile	Leu	Leu	Thr	Cys	Val	Val	865	870	875
His	Pro	Ala	Leu	Val	Phe	Asp	Ile	Thr	Lys	Leu	Leu	Leu	Ala	Ile	Phe	885	890	895
Gly	Pro	Leu	Trp	Ile	Leu	Gln	Ala	Ser	Leu	Leu	Lys	Val	Pro	Tyr	Phe	900	905	910
Val	Arg	Val	Gln	Gly	Leu	Leu	Arg	Ile	Cys	Ala	Leu	Ala	Arg	Lys	Ile	915	920	925
Ala	Gly	Gly	His	Tyr	Val	Gln	Met	Ala	Ile	Ile	Lys	Leu	Gly	Ala	Leu	930	935	940
Thr	Gly	Thr	Cys	Val	Tyr	Asn	His	Leu	Ala	Pro	Leu	Arg	Asp	Trp	Ala	945	950	955
His	Asn	Gly	Leu	Arg	Asp	Leu	Ala	Val	Ala	Val	Glu	Pro	Val	Val	Phe	965	970	975
Ser	Arg	Met	Glu	Thr	Lys	Leu	Ile	Thr	Trp	Gly	Ala	Asp	Thr	Ala	Ala	980	985	990
Cys	Gly	Asp	Ile	Ile	Asn	Gly	Leu	Pro	Val	Ser	Ala	Arg	Arg	Gly	Gln	995	1000	1005
Glu	Ile	Leu	Leu	Gly	Pro	Ala	Asp	Gly	Met	Val	Ser	Lys	Gly	Trp	Arg	1010	1015	1020
Leu	Leu	Ala	Pro	Ile	Thr	Ala	Tyr	Ala	Gln	Gln	Thr	Arg	Gly	Leu	Leu	1025	1030	1035
Gly	Cys	Ile	Ile	Thr	Ser	Leu	Thr	Gly	Arg	Asp	Lys	Asn	Gln	Val	Glu			

Ser	Glu	Pro	Ala	Pro	Ser	Gly	Cys	Pro	Pro	Asp	Ser	Asp	Val	Glu	Ser		
2370						2375				2380							
Tyr	Ser	Ser	Met	Pro	Pro	Leu	Glu	Gly	Glu	Pro	Gly	Asp	Pro	Asp	Leu		
2385				2390						2395					2400		
Ser	Asp	Gly	Ser	Trp	Ser	Thr	Val	Ser	Ser	Gly	Ala	Asp	Thr	Glu	Asp		
				2405						2410				2415			
Val	Val	Cys	Cys	Ser	Met	Ser	Tyr	Ser	Trp	Thr	Gly	Ala	Leu	Val	Thr		
				2420					2425				2430				
Pro	Cys	Ala	Ala	Glu	Glu	Gln	Lys	Leu	Pro	Ile	Asn	Ala	Leu	Ser	Asn		
				2435				2440				2445					
Ser	Leu	Leu	Arg	His	His	Asn	Leu	Val	Tyr	Ser	Thr	Thr	Ser	Arg	Ser		
				2450				2455				2460					
Ala	Cys	Gln	Arg	Lys	Lys	Lys	Val	Thr	Phe	Asp	Arg	Leu	Gln	Val	Leu		
2465					2470					2475					2480		
Asp	Ser	His	Tyr	Gln	Asp	Val	Leu	Lys	Glu	Val	Lys	Ala	Ala	Ala	Ser		
				2485					2490						2495		
Lys	Val	Lys	Ala	Asn	Leu	Leu	Ser	Val	Glu	Glu	Ala	Cys	Ser	Leu	Ala		
				2500					2505				2510				
Pro	Pro	His	Ser	Ala	Lys	Ser	Lys	Phe	Gly	Tyr	Gly	Ala	Lys	Asp	Val		
				2515				2520				2525					
Arg	Cys	His	Ala	Arg	Lys	Ala	Val	Ala	His	Ile	Asn	Ser	Val	Trp	Lys		
				2530				2535				2540					
Asp	Leu	Leu	Glu	Asp	Ser	Val	Thr	Pro	Ile	Asp	Thr	Thr	Ile	Met	Ala		
2545					2550					2555					2560		
Lys	Asn	Glu	Val	Phe	Cys	Val	Gln	Pro	Glu	Lys	Gly	Gly	Arg	Lys	Pro		
				2565					2570						2575		
Ala	Arg	Leu	Ile	Val	Phe	Pro	Asp	Leu	Gly	Val	Arg	Val	Cys	Glu	Lys		
				2580				2585					2590				
Met	Ala	Leu	Tyr	Asp	Val	Val	Ser	Lys	Leu	Pro	Leu	Ala	Val	Met	Gly		
				2595				2600				2605					
Ser	Ser	Tyr	Gly	Phe	Gln	Tyr	Ser	Pro	Gly	Gln	Arg	Val	Glu	Phe	Leu		
				2610				2615				2620					
Val	Gln	Ala	Trp	Lys	Ser	Lys	Lys	Thr	Pro	Met	Gly	Leu	Ser	Tyr	Asp		
2625					2630					2635					2640		
Thr	Arg	Cys	Phe	Asp	Ser	Thr	Val	Thr	Glu	Ser	Asp	Ile	Arg	Thr	Glu		
				2645					2650					2655			
Glu	Ala	Ile	Tyr	Gln	Cys	Cys	Asp	Leu	Asp	Pro	Gln	Ala	Arg	Val	Ala		
				2660				2665					2670				
Ile	Lys	Ser	Leu	Thr	Glu	Arg	Leu	Tyr	Val	Gly	Gly	Pro	Leu	Thr	Asn		
				2675				2680				2685					
Ser	Arg	Gly	Glu	Asn	Cys	Gly	Tyr	Arg	Arg	Cys	Arg	Ala	Ser	Arg	Val		
				2690				2695				2700					
Leu	Thr	Thr	Ser	Cys	Gly	Asn	Thr	Leu	Thr	Arg	Tyr	Ile	Lys	Ala	Arg		
2705					2710					2715					2720		
Ala	Ala	Cys	Arg	Ala	Ala	Gly	Leu	Gln	Asp	Cys	Thr	Met	Leu	Val	Cys		
				2725					2730					2735			
Gly	Asp	Asp	Leu	Val	Val	Ile	Cys	Glu	Ser	Ala	Gly	Val	Gln	Glu	Asp		
				2740				2745					2750				
Ala	Ala	Ser	Leu	Arg	Ala	Phe	Thr	Glu	Ala	Met	Thr	Arg	Tyr	Ser	Ala		
				2755				2760				2765					
Pro	Pro	Gly	Asp	Pro	Pro	Gln	Pro	Glu	Tyr	Asp	Leu	Glu	Leu	Ile	Thr		
				2770				2775				2780					
Ser	Cys	Ser	Ser	Asn	Val	Ser	Val	Ala	His	Asp	Gly	Ala	Gly	Lys	Arg		
2785					2790					2795					2800		
Val	Tyr	Tyr	Leu	Thr	Arg	Asp	Pro	Thr	Thr	Pro	Leu	Ala	Arg	Ala	Ala		

130		135		140
Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp				
145		150		155
Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile				160
	165		170	175
Phe Leu Leu Ala Leu Leu				
	180			

<210> 3
 <211> 197
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hepatitis C virus E1 protein sequence

<400> 3

Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Gln Val Arg Asn Ser Ser				
1		5		10
Gly Leu Tyr His Val Thr Asn Asp Cys Pro Asn Ser Ser Val Val Tyr				
	20		25	30
Glu Ala Ala Asp Ala Ile Leu His Thr Pro Gly Cys Val Pro Cys Val				
	35		40	45
Arg Glu Gly Asn Ala Ser Arg Cys Trp Val Ala Val Thr Pro Thr Val				
	50		55	60
Ala Thr Arg Asp Gly Lys Leu Pro Thr Thr Gln Leu Arg Arg His Ile				
	65		70	75
Asp Leu Leu Val Gly Ser Ala Thr Leu Cys Ser Ala Leu Tyr Val Gly				
	85		90	95
Asp Leu Cys Gly Ser Val Phe Leu Val Gly Gln Leu Phe Thr Phe Ser				
	100		105	110
Pro Arg His His Trp Thr Thr Gln Asp Cys Asn Cys Ser Ile Tyr Pro				
	115		120	125
Gly His Ile Thr Gly His Arg Met Ala Trp Asn Met Met Met Asn Trp				
	130		135	140
Ser Pro Thr Ala Ala Leu Val Val Ala Gln Leu Leu Arg Ile Pro Gln				
	145		150	155
Ala Ile Met Asp Met Ile Ala Gly Ala His Trp Gly Val Leu Ala Gly				
	165		170	175
Ile Lys Tyr Phe Ser Met Val Gly Asn Trp Ala Lys Val Leu Val Val				
	180		185	190
Leu Leu Leu Phe Ala				
	195			

<210> 4
 <211> 350
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hepatitis C virus E2 protein sequence

<400> 4

Gly	Val	Asp	Ala	Glu	Thr	His	Val	Thr	Gly	Gly	Asn	Ala	Gly	Arg	Thr	1	5	10	15
Thr	Ala	Gly	Leu	Val	Gly	Leu	Leu	Thr	Pro	Gly	Ala	Lys	Gln	Asn	Ile	20	25	30	
Gln	Leu	Ile	Asn	Thr	Asn	Gly	Ser	Trp	His	Ile	Asn	Ser	Thr	Ala	Leu	35	40	45	
Asn	Cys	Asn	Glu	Ser	Leu	Asn	Thr	Gly	Trp	Leu	Ala	Gly	Leu	Phe	Tyr	50	55	60	
Gln	His	Lys	Phe	Asn	Ser	Ser	Gly	Cys	Pro	Glu	Arg	Leu	Ala	Ser	Cys	65	70	75	80
Arg	Arg	Leu	Thr	Asp	Phe	Ala	Gln	Gly	Trp	Gly	Pro	Ile	Ser	Tyr	Ala	85	90	95	
Asn	Gly	Ser	Gly	Leu	Asp	Glu	Arg	Pro	Tyr	Cys	Trp	His	Tyr	Pro	Pro	100	105	110	
Arg	Pro	Cys	Gly	Ile	Val	Pro	Ala	Lys	Ser	Val	Cys	Gly	Pro	Val	Tyr	115	120	125	
Cys	Phe	Thr	Pro	Ser	Pro	Val	Val	Val	Gly	Thr	Thr	Asp	Arg	Ser	Gly	130	135	140	
Ala	Pro	Thr	Tyr	Ser	Trp	Gly	Ala	Asn	Asp	Thr	Asp	Val	Phe	Val	Leu	145	150	155	160
Asn	Asn	Thr	Arg	Pro	Pro	Leu	Gly	Asn	Trp	Phe	Gly	Cys	Thr	Trp	Met	165	170	175	
Asn	Ser	Thr	Gly	Phe	Thr	Lys	Val	Cys	Gly	Ala	Pro	Pro	Cys	Val	Ile	180	185	190	
Gly	Gly	Val	Gly	Asn	Asn	Thr	Leu	Leu	Cys	Pro	Thr	Asp	Cys	Phe	Arg	195	200	205	
Lys	Tyr	Pro	Glu	Ala	Thr	Tyr	Ser	Arg	Cys	Gly	Ser	Gly	Pro	Arg	Ile	210	215	220	
Thr	Pro	Arg	Cys	Met	Val	Asp	Tyr	Pro	Tyr	Arg	Leu	Trp	His	Tyr	Pro	225	230	235	240
Cys	Thr	Ile	Asn	Tyr	Thr	Ile	Phe	Lys	Val	Arg	Met	Tyr	Val	Gly	Gly	245	250	255	
Val	Glu	His	Arg	Leu	Glu	Ala	Ala	Cys	Asn	Trp	Thr	Arg	Gly	Glu	Arg	260	265	270	
Cys	Asp	Leu	Glu	Asp	Arg	Asp	Arg	Ser	Glu	Leu	Ser	Pro	Leu	Leu	Leu	275	280	285	
Ser	Thr	Thr	Gln	Trp	Gln	Val	Leu	Pro	Cys	Ser	Phe	Thr	Thr	Leu	Pro	290	295	300	
Ala	Leu	Ser	Thr	Gly	Leu	Ile	His	Leu	His	Gln	Asn	Ile	Val	Asp	Val	305	310	315	320
Gln	Tyr	Leu	Tyr	Gly	Val	Gly	Ser	Ser	Ile	Ala	Ser	Trp	Ala	Ile	Lys	325	330	335	
Trp	Glu	Tyr	Val	Val	Leu	Leu	Phe	Leu	Leu	Leu	Ala	Asp	Ala	340	345	350			

<210> 5

<211> 315

<212> PRT

<213> Artificial Sequence

<220>

<223> Hepatitis C virus NS2 protein sequence

<400> 5

Arg	Val	Cys	Ser	Cys	Leu	Trp	Met	Met	Leu	Leu	Ile	Ser	Gln	Ala	Glu	1	5	10	15
Ala	Ala	Leu	Glu	Asn	Leu	Val	Ile	Leu	Asn	Ala	Ala	Ser	Leu	Ala	Gly	20	25	30	
Thr	His	Gly	Leu	Val	Ser	Phe	Leu	Val	Phe	Phe	Cys	Phe	Ala	Trp	Tyr	35	40	45	
Leu	Lys	Gly	Arg	Trp	Val	Pro	Gly	Ala	Val	Tyr	Ala	Leu	Tyr	Gly	Met	50	55	60	
Trp	Pro	Leu	Leu	Leu	Leu	Leu	Leu	Ala	Leu	Pro	Gln	Arg	Ala	Tyr	Ala	65	70	75	80
Leu	Asp	Thr	Glu	Val	Ala	Ala	Ser	Cys	Gly	Gly	Val	Val	Leu	Val	Gly	85	90	95	
Leu	Met	Ala	Leu	Thr	Leu	Ser	Pro	Tyr	Tyr	Lys	Arg	Tyr	Ile	Ser	Trp	100	105	110	
Cys	Met	Trp	Trp	Leu	Gln	Tyr	Phe	Leu	Thr	Arg	Val	Glu	Ala	Gln	Leu	115	120	125	
His	Val	Trp	Val	Pro	Pro	Leu	Asn	Val	Arg	Gly	Gly	Arg	Asp	Ala	Val	130	135	140	
Ile	Leu	Leu	Thr	Cys	Val	Val	His	Pro	Ala	Leu	Val	Phe	Asp	Ile	Thr	145	150	155	160
Lys	Leu	Leu	Leu	Ala	Ile	Phe	Gly	Pro	Leu	Trp	Ile	Leu	Gln	Ala	Ser	165	170	175	
Leu	Leu	Lys	Val	Pro	Tyr	Phe	Val	Arg	Val	Gln	Gly	Leu	Leu	Arg	Ile	180	185	190	
Cys	Ala	Leu	Ala	Arg	Lys	Ile	Ala	Gly	Gly	His	Tyr	Val	Gln	Met	Ala	195	200	205	
Ile	Ile	Lys	Leu	Gly	Ala	Leu	Thr	Gly	Thr	Cys	Val	Tyr	Asn	His	Leu	210	215	220	
Ala	Pro	Leu	Arg	Asp	Trp	Ala	His	Asn	Gly	Leu	Arg	Asp	Leu	Ala	Val	225	230	235	240
Ala	Val	Glu	Pro	Val	Val	Phe	Ser	Arg	Met	Glu	Thr	Lys	Leu	Ile	Thr	245	250	255	
Trp	Gly	Ala	Asp	Thr	Ala	Ala	Cys	Gly	Asp	Ile	Ile	Asn	Gly	Leu	Pro	260	265	270	
Val	Ser	Ala	Arg	Arg	Gly	Gln	Glu	Ile	Leu	Leu	Gly	Pro	Ala	Asp	Gly	275	280	285	
Met	Val	Ser	Lys	Gly	Trp	Arg	Leu	Leu	Ala	Pro	Ile	Thr	Ala	Tyr	Ala	290	295	300	
Gln	Gln	Thr	Arg	Gly	Leu	Leu	Gly	Cys	Ile	Ile	305	310	315						

<210> 6

<211> 613

<212> PRT

<213> Artificial Sequence

<220>

<223> Hepatitis C virus NS3 protein sequence

<400> 6

Thr	Ser	Leu	Thr	Gly	Arg	Asp	Lys	Asn	Gln	Val	Glu	Gly	Glu	Val	Gln	1	5	10	15
Ile	Val	Ser	Thr	Ala	Thr	Gln	Thr	Phe	Leu	Ala	Thr	Cys	Ile	Asn	Gly	20	25	30	

Val	Cys	Trp	Thr	Val	Tyr	His	Gly	Ala	Gly	Thr	Arg	Thr	Ile	Ala	Ser			
		35					40					45						
Pro	Lys	Gly	Pro	Val	Ile	Gln	Thr	Tyr	Thr	Asn	Val	Asp	Gln	Asp	Leu			
		50				55				60								
Val	Gly	Trp	Pro	Ala	Pro	Gln	Gly	Ser	Arg	Ser	Leu	Thr	Pro	Cys	Thr			
65					70					75					80			
Cys	Gly	Ser	Ser	Asp	Leu	Tyr	Leu	Val	Thr	Arg	His	Ala	Asp	Val	Ile			
				85					90					95				
Pro	Val	Arg	Arg	Arg	Gly	Asp	Ser	Arg	Gly	Ser	Leu	Leu	Ser	Pro	Arg			
				100				105						110				
Pro	Ile	Ser	Tyr	Leu	Lys	Gly	Ser	Ser	Gly	Gly	Pro	Leu	Leu	Cys	Pro			
		115				120								125				
Thr	Gly	His	Ala	Val	Gly	Leu	Phe	Arg	Ala	Ala	Val	Cys	Thr	Arg	Gly			
		130				135					140							
Val	Ala	Lys	Ala	Val	Asp	Phe	Ile	Pro	Val	Glu	Asn	Leu	Glu	Thr	Thr			
145					150					155					160			
Met	Arg	Ser	Pro	Val	Phe	Thr	Asp	Asn	Ser	Ser	Pro	Pro	Ala	Val	Pro			
				165				170						175				
Gln	Ser	Phe	Gln	Val	Ala	His	Leu	His	Ala	Pro	Thr	Gly	Ser	Gly	Lys			
			180					185						190				
Ser	Thr	Lys	Val	Pro	Ala	Ala	Tyr	Ala	Ala	Lys	Gly	Tyr	Lys	Val	Leu			
		195					200							205				
Val	Leu	Asn	Pro	Ser	Val	Ala	Ala	Thr	Leu	Gly	Phe	Gly	Ala	Tyr	Met			
		210				215						220						
Ser	Lys	Ala	His	Gly	Val	Asp	Pro	Asn	Ile	Arg	Thr	Gly	Val	Arg	Thr			
225					230					235					240			
Ile	Thr	Thr	Gly	Ser	Pro	Ile	Thr	Tyr	Ser	Thr	Tyr	Gly	Lys	Phe	Leu			
				245				250						255				
Ala	Asp	Ala	Gly	Cys	Ser	Gly	Gly	Ala	Tyr	Asp	Ile	Ile	Ile	Cys	Asp			
			260					265						270				
Glu	Cys	His	Ser	Thr	Asp	Ala	Thr	Ser	Ile	Ser	Gly	Ile	Gly	Thr	Val			
		275					280					285						
Leu	Asp	Gln	Ala	Glu	Thr	Ala	Gly	Ala	Arg	Leu	Val	Val	Leu	Ala	Thr			
		290				295					300							
Ala	Thr	Pro	Pro	Gly	Ser	Val	Thr	Val	Ser	His	Pro	Asn	Ile	Glu	Glu			
305					310					315					320			
Val	Ala	Leu	Ser	Thr	Thr	Gly	Glu	Ile	Pro	Phe	Tyr	Gly	Lys	Ala	Ile			
				325					330					335				
Pro	Leu	Glu	Val	Ile	Lys	Gly	Gly	Arg	His	Leu	Ile	Phe	Cys	His	Ser			
			340					345						350				
Lys	Lys	Lys	Cys	Asp	Glu	Leu	Ala	Ala	Lys	Leu	Val	Ala	Leu	Gly	Ile			
		355					360							365				
Asn	Ala	Val	Ala	Tyr	Tyr	Arg	Gly	Leu	Asp	Val	Ser	Val	Ile	Pro	Thr			
		370				375								380				
Ser	Gly	Asp	Val	Val	Val	Val	Ser	Thr	Asp	Ala	Leu	Met	Thr	Gly	Phe			
385					390					395					400			
Thr	Gly	Asp	Phe	Asp	Ser	Val	Ile	Asp	Cys	Asn	Thr	Cys	Val	Thr	Gln			
				405					410					415				
Thr	Val	Asp	Phe	Ser	Leu	Asp	Pro	Thr	Phe	Thr	Ile	Glu	Thr	Thr	Thr			
			420					425						430				
Leu	Pro	Gln	Asp	Ala	Val	Ser	Arg	Thr	Gln	Arg	Arg	Gly	Arg	Thr	Gly			
		435					440					445						
Arg	Gly	Lys	Pro	Gly	Ile	Tyr	Arg	Phe	Val	Ala	Pro	Gly	Glu	Arg	Pro			
		450				455					460							
Ser	Gly	Met	Phe	Asp	Ser	Ser	Val	Leu	Cys	Glu	Cys	Tyr	Asp	Ala	Gly			

465		470		475		480									
Cys	Ala	Trp	Tyr	Glu	Leu	Thr	Pro	Ala	Glu	Thr	Thr	Val	Arg	Leu	Arg
		485		490		495									
Ala	Tyr	Met	Asn	Thr	Pro	Gly	Leu	Pro	Val	Cys	Gln	Asp	His	Leu	Gly
		500		505		510									
Phe	Trp	Glu	Gly	Val	Phe	Thr	Gly	Leu	Thr	His	Ile	Asp	Ala	His	Phe
		515		520		525									
Leu	Ser	Gln	Thr	Lys	Gln	Ser	Gly	Glu	Asn	Phe	Pro	Tyr	Leu	Val	Ala
		530		535		540									
Tyr	Gln	Ala	Thr	Val	Cys	Ala	Arg	Ala	Gln	Ala	Pro	Pro	Pro	Ser	Trp
		545		550		555									
Asp	Gln	Met	Arg	Lys	Cys	Leu	Ile	Arg	Leu	Lys	Pro	Thr	Leu	His	Gly
		565		570		575									
Pro	Thr	Pro	Leu	Leu	Tyr	Arg	Leu	Gly	Ala	Val	Gln	Asn	Glu	Val	Thr
		580		585		590									
Leu	Thr	His	Pro	Ile	Thr	Lys	Tyr	Ile	Met	Thr	Cys	Met	Ser	Ala	Asp
		595		600		605									
Leu	Glu	Val	Val	Thr											
		610													

<210> 7
 <211> 54
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hepatitis C virus NS4A protein sequence

<400> 7
Ser Thr Trp Val Leu Val Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr
1 5 10 15
Cys Leu Ser Thr Gly Cys Val Val Ile Val Gly Arg Ile Val Leu Ser
20 25 30
Gly Lys Pro Ala Ile Ile Pro Asp Arg Glu Val Leu Tyr Gln Glu Phe
35 40 45
Asp Glu Met Glu Glu Cys
50

<210> 8
 <211> 260
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hepatitis C virus NS4B protein sequence

<400> 8
Ser Gln His Leu Pro Tyr Ile Glu Gln Gly Met Met Leu Ala Glu Gln
1 5 10 15
Phe Lys Gln Lys Ala Leu Gly Leu Leu Gln Thr Ala Ser Arg His Ala
20 25 30
Glu Val Ile Thr Pro Ala Val Gln Thr Asn Trp Gln Lys Leu Glu Val
35 40 45

Phe	Trp	Ala	Lys	His	Met	Trp	Asn	Phe	Ile	Ser	Gly	Ile	Gln	Tyr	Leu
50						55					60				
Ala	Gly	Leu	Ser	Thr	Leu	Pro	Gly	Asn	Pro	Ala	Ile	Ala	Ser	Leu	Met
65						70				75					80
Ala	Phe	Thr	Ala	Ala	Val	Thr	Ser	Pro	Leu	Thr	Thr	Gly	Gln	Thr	Leu
				85					90					95	
Leu	Phe	Asn	Ile	Leu	Gly	Gly	Trp	Val	Ala	Ala	Gln	Leu	Ala	Ala	Pro
			100					105					110		
Gly	Ala	Ala	Thr	Ala	Phe	Val	Gly	Ala	Gly	Leu	Ala	Gly	Ala	Ala	Leu
			115					120					125		
Asp	Ser	Val	Gly	Leu	Gly	Lys	Val	Leu	Val	Asp	Ile	Leu	Ala	Gly	Tyr
			130				135					140			
Gly	Ala	Gly	Val	Ala	Gly	Ala	Leu	Val	Ala	Phe	Lys	Ile	Met	Ser	Gly
145						150				155					160
Glu	Val	Pro	Ser	Thr	Glu	Asp	Leu	Val	Asn	Leu	Leu	Pro	Ala	Ile	Leu
					165				170					175	
Ser	Pro	Gly	Ala	Leu	Ala	Val	Gly	Val	Val	Phe	Ala	Ser	Ile	Leu	Arg
			180					185					190		
Arg	Arg	Val	Gly	Pro	Gly	Glu	Gly	Ala	Val	Gln	Trp	Met	Asn	Arg	Leu
			195					200					205		
Ile	Ala	Phe	Ala	Ser	Arg	Gly	Asn	His	Val	Ser	Pro	Thr	His	Tyr	Val
			210					215					220		
Pro	Glu	Ser	Asp	Ala	Ala	Ala	Arg	Val	Thr	Ala	Ile	Leu	Ser	Ser	Leu
225						230				235					240
Thr	Val	Thr	Gln	Leu	Leu	Arg	Arg	Leu	His	Gln	Trp	Ile	Ser	Ser	Glu
				245					250					255	
Cys	Thr	Thr	Pro												
			260												

<210> 9

<211> 1040

<212> PRT

<213> Artificial Sequence

<220>

<223> Hepatitis C virus NS5A/B protein sequence

<400> 9

Cys	Ser	Gly	Ser	Trp	Leu	Arg	Asp	Ile	Trp	Asp	Trp	Ile	Cys	Glu	Val
1				5					10				15		
Leu	Ser	Asp	Phe	Lys	Thr	Trp	Leu	Lys	Ala	Lys	Leu	Met	Pro	Gln	Leu
			20					25				30			
Pro	Gly	Ile	Pro	Phe	Val	Ser	Cys	Gln	Arg	Gly	Tyr	Arg	Gly	Val	Trp
			35				40					45			
Arg	Gly	Asp	Gly	Ile	Met	His	Thr	Arg	Cys	His	Cys	Gly	Ala	Glu	Ile
			50			55					60				
Thr	Gly	His	Val	Lys	Asn	Gly	Thr	Met	Arg	Ile	Val	Gly	Pro	Arg	Thr
65					70				75					80	
Cys	Lys	Asn	Met	Trp	Ser	Gly	Thr	Phe	Phe	Ile	Asn	Ala	Tyr	Thr	Thr
			85					90					95		
Gly	Pro	Cys	Thr	Pro	Leu	Pro	Ala	Pro	Asn	Tyr	Lys	Phe	Ala	Leu	Trp
			100					105				110			
Arg	Val	Ser	Ala	Glu	Glu	Tyr	Val	Glu	Ile	Arg	Arg	Val	Gly	Asp	Phe
			115					120					125		

His	Tyr	Val	Ser	Gly	Met	Thr	Thr	Asp	Asn	Leu	Lys	Cys	Pro	Cys	Gln
130						135					140				
Ile	Pro	Ser	Pro	Glu	Phe	Phe	Thr	Glu	Leu	Asp	Gly	Val	Arg	Leu	His
145					150					155					160
Arg	Phe	Ala	Pro	Pro	Cys	Lys	Pro	Leu	Leu	Arg	Glu	Glu	Val	Ser	Phe
					165				170					175	
Arg	Val	Gly	Leu	His	Glu	Tyr	Pro	Val	Gly	Ser	Gln	Leu	Pro	Cys	Glu
					180				185				190		
Pro	Glu	Pro	Asp	Val	Ala	Val	Leu	Thr	Ser	Met	Leu	Thr	Asp	Pro	Ser
					195				200				205		
His	Ile	Thr	Ala	Glu	Ala	Ala	Gly	Arg	Arg	Leu	Ala	Arg	Gly	Ser	Pro
210							215				220				
Pro	Ser	Met	Ala	Ser	Ser	Ser	Ala	Ser	Gln	Leu	Ser	Ala	Pro	Ser	Leu
225					230					235					240
Lys	Ala	Thr	Cys	Thr	Ala	Asn	His	Asp	Ser	Pro	Asp	Ala	Glu	Leu	Ile
					245				250					255	
Glu	Ala	Asn	Leu	Leu	Trp	Arg	Gln	Glu	Met	Gly	Gly	Asn	Ile	Thr	Arg
					260				265				270		
Val	Glu	Ser	Glu	Asn	Lys	Val	Val	Ile	Leu	Asp	Ser	Phe	Asp	Pro	Leu
					275				280				285		
Val	Ala	Glu	Glu	Asp	Glu	Arg	Glu	Val	Ser	Val	Pro	Ala	Glu	Ile	Leu
							295				300				
Arg	Lys	Ser	Arg	Arg	Phe	Ala	Pro	Ala	Leu	Pro	Val	Trp	Ala	Arg	Pro
305					310					315					320
Asp	Tyr	Asn	Pro	Leu	Leu	Val	Glu	Thr	Trp	Lys	Lys	Pro	Asp	Tyr	Glu
					325				330					335	
Pro	Pro	Val	Val	His	Gly	Cys	Pro	Leu	Pro	Pro	Pro	Arg	Ser	Pro	Pro
					340				345				350		
Val	Pro	Pro	Pro	Arg	Lys	Lys	Arg	Thr	Val	Val	Leu	Thr	Glu	Ser	Thr
					355				360				365		
Leu	Pro	Thr	Ala	Leu	Ala	Glu	Leu	Ala	Thr	Lys	Ser	Phe	Gly	Ser	Ser
						375					380				
Ser	Thr	Ser	Gly	Ile	Thr	Gly	Asp	Asn	Thr	Thr	Thr	Ser	Ser	Glu	Pro
385					390					395					400
Ala	Pro	Ser	Gly	Cys	Pro	Pro	Asp	Ser	Asp	Val	Glu	Ser	Tyr	Ser	Ser
					405				410					415	
Met	Pro	Pro	Leu	Glu	Gly	Glu	Pro	Gly	Asp	Pro	Asp	Leu	Ser	Asp	Gly
					420				425				430		
Ser	Trp	Ser	Thr	Val	Ser	Ser	Gly	Ala	Asp	Thr	Glu	Asp	Val	Val	Cys
					435				440				445		
Cys	Ser	Met	Ser	Tyr	Ser	Trp	Thr	Gly	Ala	Leu	Val	Thr	Pro	Cys	Ala
						455					460				
Ala	Glu	Glu	Gln	Lys	Leu	Pro	Ile	Asn	Ala	Leu	Ser	Asn	Ser	Leu	Leu
465					470					475					480
Arg	His	His	Asn	Leu	Val	Tyr	Ser	Thr	Thr	Ser	Arg	Ser	Ala	Cys	Gln
					485				490					495	
Arg	Lys	Lys	Lys	Val	Thr	Phe	Asp	Arg	Leu	Gln	Val	Leu	Asp	Ser	His
					500				505				510		
Tyr	Gln	Asp	Val	Leu	Lys	Glu	Val	Lys	Ala	Ala	Ala	Ser	Lys	Val	Lys
					515				520				525		
Ala	Asn	Leu	Leu	Ser	Val	Glu	Glu	Ala	Cys	Ser	Leu	Ala	Pro	Pro	His
					530				535				540		
Ser	Ala	Lys	Ser	Lys	Phe	Gly	Tyr	Gly	Ala	Lys	Asp	Val	Arg	Cys	His
545					550					555					560
Ala	Arg	Lys	Ala	Val	Ala	His	Ile	Asn	Ser	Val	Trp	Lys	Asp	Leu	Leu

Ile Tyr His Ser Val Ser His Ala Arg Pro Arg Trp Phe Trp Phe Cys
 1010 1015 1020
 Leu Leu Leu Leu Ala Ala Gly Val Gly Ile Tyr Leu Leu Pro Asn Arg
 1025 1030 1035 1040

<210> 10
 <211> 226
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hepatitis B virus S antigen (HBsAg) sequence

<400> 10
 Met Glu Asn Ile Thr Ser Gly Phe Leu Gly Pro Leu Leu Val Leu Gln
 1 5 10 15
 Ala Gly Phe Phe Leu Leu Thr Arg Ile Leu Thr Ile Pro Gln Ser Leu
 20 25 30
 Asp Ser Trp Trp Thr Ser Leu Asn Phe Leu Gly Gly Thr Thr Val Cys
 35 40 45
 Leu Gly Gln Asn Ser Gln Ser Pro Thr Ser Asn His Ser Pro Thr Ser
 50 55 60
 Cys Pro Pro Thr Cys Pro Gly Tyr Arg Trp Met Cys Leu Arg Arg Phe
 65 70 75 80
 Ile Ile Phe Leu Phe Ile Leu Leu Leu Cys Leu Ile Phe Leu Leu Val
 85 90 95
 Leu Leu Asp Tyr Gln Gly Met Leu Pro Val Cys Pro Leu Ile Pro Gly
 100 105 110
 Ser Ser Thr Thr Ser Thr Gly Pro Cys Arg Thr Cys Met Thr Thr Ala
 115 120 125
 Gln Gly Thr Ser Met Tyr Pro Ser Cys Cys Cys Thr Lys Pro Ser Asp
 130 135 140
 Gly Asn Cys Thr Cys Ile Pro Ile Pro Ser Ser Trp Ala Phe Gly Lys
 145 150 155 160
 Phe Leu Trp Glu Trp Ala Ser Ala Arg Phe Ser Trp Leu Ser Leu Leu
 165 170 175
 Val Pro Phe Val Gln Trp Phe Val Gly Leu Ser Pro Thr Val Trp Leu
 180 185 190
 Ser Val Ile Trp Met Met Trp Tyr Trp Gly Pro Ser Leu Tyr Ser Ile
 195 200 205
 Leu Ser Pro Phe Leu Pro Leu Leu Pro Ile Phe Phe Cys Leu Trp Val
 210 215 220
 Tyr Ile
 225

<210> 11
 <211> 212
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hepatitis B virus C antigen and e antigen
 (HBcAg/HBeAg) sequence

<400> 11

```
Met Gln Leu Phe His Leu Cys Leu Ile Ile Ser Cys Ser Cys Pro Thr
 1          5          10          15
Val Gln Ala Ser Lys Leu Cys Leu Gly Trp Leu Trp Gly Met Asp Ile
          20          25          30
Asp Pro Tyr Lys Glu Phe Gly Ala Thr Val Glu Leu Leu Ser Phe Leu
          35          40          45
Pro Ser Asp Phe Phe Pro Ser Val Arg Asp Leu Leu Asp Thr Ala Ser
          50          55          60
Ala Leu Tyr Arg Glu Ala Leu Glu Ser Pro Glu His Cys Ser Pro His
65          70          75          80
His Thr Ala Leu Arg Gln Ala Ile Leu Cys Trp Gly Glu Leu Met Thr
          85          90          95
Leu Ala Thr Trp Val Gly Val Asn Leu Glu Asp Pro Ala Ser Arg Asp
          100          105          110
Leu Val Val Ser Tyr Val Asn Thr Asn Met Gly Leu Lys Phe Arg Gln
          115          120          125
Leu Leu Trp Phe His Ile Ser Cys Leu Thr Phe Gly Arg Glu Thr Val
          130          135          140
Ile Glu Tyr Leu Val Ser Phe Gly Val Trp Ile Arg Thr Pro Pro Ala
145          150          155          160
Tyr Arg Pro Pro Asn Ala Pro Ile Leu Ser Thr Leu Pro Glu Thr Thr
          165          170          175
Val Val Arg Arg Arg Gly Arg Ser Pro Arg Arg Arg Thr Pro Ser Pro
          180          185          190
Arg Arg Arg Arg Ser Gln Ser Pro Arg Arg Arg Arg Ser Gln Ser Arg
          195          200          205
Glu Ser Gln Cys
          210
```

<210> 12

<211> 2227

<212> PRT

<213> Artificial Sequence

<220>

<223> Hepatitis A virus sequence

<400> 12

```
Met Asn Met Ser Lys Gln Gly Ile Phe Gln Thr Val Gly Ser Gly Leu
 1          5          10          15
Asp His Ile Leu Ser Leu Ala Asp Ile Glu Glu Glu Gln Met Ile Gln
          20          25          30
Ser Val Asp Arg Thr Ala Val Thr Gly Ala Ser Tyr Phe Thr Ser Val
          35          40          45
Asp Gln Ser Ser Val His Thr Ala Glu Val Gly Ser His Gln Ile Glu
          50          55          60
Pro Leu Lys Thr Ser Val Asp Lys Pro Gly Ser Lys Lys Thr Gln Gly
65          70          75          80
Glu Lys Phe Phe Leu Ile His Ser Ala Asp Trp Leu Thr Thr His Ala
          85          90          95
Leu Phe His Glu Val Ala Lys Leu Asp Val Val Lys Leu Leu Tyr Asn
          100          105          110
```

Glu	Gln	Phe	Ala	Val	Gln	Gly	Leu	Leu	Arg	Tyr	His	Thr	Tyr	Ala	Arg	115	120	125
Phe	Gly	Ile	Glu	Ile	Gln	Val	Gln	Ile	Asn	Pro	Thr	Pro	Phe	Gln	Gln	130	135	140
Gly	Gly	Leu	Ile	Cys	Ala	Met	Val	Pro	Gly	Asp	Gln	Ser	Tyr	Gly	Ser	145	150	155
Ile	Ala	Ser	Leu	Thr	Val	Tyr	Pro	His	Gly	Leu	Leu	Asn	Cys	Asn	Ile	165	170	175
Asn	Asn	Val	Val	Arg	Ile	Lys	Val	Pro	Phe	Ile	Tyr	Thr	Arg	Gly	Ala	180	185	190
Tyr	His	Phe	Lys	Asp	Pro	Gln	Tyr	Pro	Val	Trp	Glu	Leu	Thr	Ile	Arg	195	200	205
Val	Trp	Ser	Glu	Leu	Asn	Ile	Gly	Thr	Gly	Thr	Ser	Ala	Tyr	Thr	Ser	210	215	220
Leu	Asn	Val	Leu	Ala	Arg	Phe	Thr	Asp	Leu	Glu	Leu	His	Gly	Leu	Thr	225	230	235
Pro	Leu	Ser	Thr	Gln	Met	Met	Arg	Asn	Glu	Phe	Arg	Val	Ser	Thr	Thr	245	250	255
Glu	Asn	Val	Val	Asn	Leu	Ser	Asn	Tyr	Glu	Asp	Ala	Arg	Ala	Lys	Met	260	265	270
Ser	Phe	Ala	Leu	Asp	Gln	Glu	Asp	Trp	Lys	Ser	Asp	Pro	Ser	Gln	Gly	275	280	285
Gly	Gly	Ile	Lys	Ile	Thr	His	Phe	Thr	Thr	Trp	Thr	Ser	Ile	Pro	Thr	290	295	300
Leu	Ala	Ala	Gln	Phe	Pro	Phe	Asn	Ala	Ser	Asp	Ser	Val	Gly	Gln	Gln	305	310	315
Ile	Lys	Val	Ile	Pro	Val	Asp	Pro	Tyr	Phe	Phe	Gln	Met	Thr	Asn	Thr	325	330	335
Asn	Pro	Asp	Gln	Lys	Cys	Ile	Thr	Ala	Leu	Ala	Ser	Ile	Cys	Gln	Met	340	345	350
Phe	Cys	Phe	Trp	Arg	Gly	Asp	Leu	Val	Phe	Asp	Phe	Gln	Val	Phe	Pro	355	360	365
Thr	Lys	Tyr	His	Ser	Gly	Arg	Leu	Leu	Phe	Cys	Phe	Val	Pro	Gly	Asn	370	375	380
Glu	Leu	Ile	Asp	Val	Thr	Gly	Ile	Thr	Leu	Lys	Gln	Ala	Thr	Thr	Ala	385	390	395
Pro	Cys	Ala	Val	Met	Asp	Ile	Thr	Gly	Val	Gln	Ser	Thr	Leu	Arg	Phe	405	410	415
Arg	Val	Pro	Trp	Ile	Ser	Asp	Thr	Pro	Tyr	Arg	Val	Asn	Arg	Tyr	Thr	420	425	430
Lys	Ser	Ala	His	Gln	Lys	Gly	Glu	Tyr	Thr	Ala	Ile	Gly	Lys	Leu	Ile	435	440	445
Val	Tyr	Cys	Tyr	Asn	Arg	Leu	Thr	Ser	Pro	Ser	Asn	Val	Ala	Ser	His	450	455	460
Val	Arg	Val	Asn	Val	Tyr	Leu	Ser	Ala	Ile	Asn	Leu	Glu	Cys	Phe	Ala	465	470	475
Pro	Leu	Tyr	His	Ala	Met	Asp	Val	Thr	Thr	Gln	Val	Gly	Asp	Asp	Ser	485	490	495
Gly	Gly	Phe	Ser	Thr	Thr	Val	Ser	Thr	Glu	Gln	Asn	Val	Pro	Asp	Pro	500	505	510
Gln	Val	Gly	Ile	Thr	Thr	Met	Arg	Asp	Leu	Lys	Gly	Lys	Ala	Asn	Arg	515	520	525
Gly	Lys	Met	Asp	Val	Ser	Gly	Val	Gln	Ala	Pro	Arg	Gly	Ser	Tyr	Gln	530	535	540
Gln	Gln	Leu	Asn	Asp	Pro	Val	Leu	Ala	Lys	Lys	Val	Pro	Glu	Thr	Phe			

545	Pro	Glu	Leu	Lys	Pro	Gly	Glu	Ser	Arg	His	Thr	Ser	Asp	His	Met	Ser
					565					570					575	
Ile	Tyr	Lys	Phe	Met	Gly	Arg	Ser	His	Phe	Leu	Cys	Thr	Phe	Thr	Phe	
					580				585					590		
Asn	Ser	Asn	Asn	Lys	Glu	Tyr	Thr	Phe	Pro	Ile	Thr	Leu	Ser	Ser	Thr	
					595				600					605		
Ser	Asn	Pro	Pro	His	Gly	Leu	Pro	Ser	Thr	Leu	Arg	Trp	Phe	Phe	Asn	
					610				615					620		
Leu	Phe	Gln	Leu	Tyr	Arg	Gly	Pro	Leu	Asp	Leu	Thr	Ile	Ile	Ile	Thr	
625					630					635					640	
Gly	Ala	Thr	Asp	Val	Asp	Gly	Met	Ala	Trp	Phe	Thr	Pro	Val	Gly	Leu	
					645					650					655	
Ala	Val	Asp	Pro	Trp	Val	Glu	Lys	Glu	Ser	Ala	Leu	Ser	Ile	Asp	Tyr	
					660					665					670	
Lys	Thr	Ala	Leu	Gly	Ala	Val	Arg	Phe	Asn	Thr	Arg	Arg	Thr	Gly	Asn	
					675					680					685	
Ile	Gln	Ile	Arg	Leu	Pro	Trp	Tyr	Ser	Tyr	Leu	Tyr	Ala	Val	Ser	Gly	
					690					695					700	
Ala	Leu	Asp	Gly	Leu	Gly	Asp	Lys	Thr	Asp	Ser	Thr	Phe	Gly	Leu	Phe	
705					710					715					720	
Leu	Phe	Glu	Ile	Ala	Asn	Tyr	Asn	His	Ser	Asp	Glu	Tyr	Leu	Ser	Phe	
					725					730					735	
Ser	Cys	Tyr	Leu	Ser	Val	Thr	Glu	Gln	Ser	Glu	Phe	Tyr	Phe	Pro	Arg	
					740					745					750	
Ala	Pro	Leu	Asn	Ser	Asn	Ala	Met	Leu	Ser	Thr	Glu	Ser	Met	Met	Ser	
					755					760					765	
Arg	Ile	Ala	Ala	Gly	Asp	Leu	Glu	Ser	Ser	Val	Asp	Asp	Pro	Arg	Ser	
					770					775					780	
Glu	Glu	Asp	Arg	Arg	Phe	Glu	Ser	His	Ile	Glu	Cys	Arg	Lys	Pro	Tyr	
785					790					795					800	
Lys	Glu	Leu	Arg	Leu	Glu	Val	Gly	Lys	Gln	Arg	Leu	Lys	Tyr	Ala	Gln	
					805					810					815	
Glu	Glu	Leu	Ser	Asn	Glu	Val	Leu	Pro	Pro	Pro	Arg	Lys	Met	Lys	Gly	
					820					825					830	
Leu	Phe	Ser	Gln	Ala	Lys	Ile	Ser	Leu	Phe	Tyr	Thr	Glu	Glu	His	Glu	
					835					840					845	
Ile	Met	Lys	Phe	Ser	Trp	Arg	Gly	Val	Thr	Ala	Asp	Thr	Arg	Ala	Leu	
					850					855					860	
Arg	Arg	Phe	Gly	Phe	Ser	Leu	Ala	Ala	Gly	Arg	Ser	Val	Trp	Thr	Leu	
865					870					875					880	
Glu	Met	Asp	Ala	Gly	Val	Leu	Thr	Gly	Arg	Leu	Ile	Arg	Leu	Asn	Asp	
					885					890					895	
Glu	Lys	Trp	Thr	Glu	Met	Lys	Asp	Asp	Lys	Ile	Val	Ser	Leu	Ile	Glu	
					900					905					910	
Lys	Phe	Thr	Ser	Asn	Lys	Tyr	Trp	Ser	Lys	Val	Asn	Phe	Pro	His	Gly	
					915					920					925	
Met	Leu	Asp	Leu	Glu	Glu	Ile	Ala	Ala	Asn	Ser	Lys	Asp	Phe	Pro	Asn	
					930					935					940	
Met	Ser	Glu	Thr	Asp	Leu	Cys	Phe	Leu	Leu	His	Trp	Leu	Asn	Pro	Lys	
945					950					955					960	
Lys	Ile	Asn	Leu	Ala	Asp	Arg	Met	Leu	Gly	Leu	Ser	Gly	Val	Gln	Glu	
					965					970					975	
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Gly Val Leu Val Gly Gly Trp Phe Val Tyr Lys His Phe Ser Arg Lys						
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Glu Glu Glu Pro Ile Pro Ala Glu Gly Val Tyr His Gly Val Thr Lys						
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Pro Lys Gln Val Ile Lys Leu Asp Ala Asp Pro Val Glu Ser Gln Ser						
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Thr Leu Glu Ile Ala Gly Leu Val Arg Lys Asn Leu Val Gln Phe Gly						
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Val Lys Asp Asp Trp Leu Leu Val Pro Ser His Ala Tyr Lys Phe Glu						
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Lys Asp Tyr Glu Met Met Glu Phe Tyr Phe Asn Arg Gly Gly Thr Tyr						
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<213> Artificial Sequence

<220>

<223> Hepatitis B virus sequence

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<211> 7478

<212> DNA

<213> Artificial Sequence

<220>

<223> Hepatitis A virus sequence

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<223> Hepatitis C virus NS3/4A peptide

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Cys Ile Ile Thr Ser Leu Thr Gly Arg Asp Lys Asn Gln Val Glu Gly
20 25 30
Glu Val Gln Ile Val Ser Thr Ala Ala Gln Thr Phe Leu Ala Thr Cys
35 40 45
Ile Asn Gly Val Cys Trp Thr Val Tyr His Gly Ala Gly Thr Arg Thr
50 55 60
Ile Ala Ser Pro Lys Gly Pro Val Ile Gln Met Tyr Thr Asn Val Asp
65 70 75 80
Gln Asp Leu Val Gly Trp Pro Ala Pro Gln Gly Ala Arg Ser Leu Thr

His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr His Ile Asp
 530 535 540
 Ala His Phe Leu Ser Gln Thr Lys Gln Ser Gly Glu Asn Leu Pro Tyr
 545 550 555 560
 Leu Val Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala Gln Ala Pro Pro
 565 570 575
 Pro Ser Trp Asp Gln Met Trp Lys Cys Leu Ile Arg Leu Lys Pro Thr
 580 585 590
 Leu His Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Ala Val Gln Asn
 595 600 605
 Glu Val Thr Leu Thr His Pro Val Thr Lys Tyr Ile Met Thr Cys Met
 610 615 620
 Ser Ala Asp Leu Glu Val Val Thr
 625 630

<210> 30
 <211> 54
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hepatitis C virus NS4A peptide

<400> 30
 Ser Thr Trp Val Leu Val Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr
 1 5 10 15
 Cys Leu Ser Thr Gly Cys Val Val Ile Val Gly Arg Ile Val Leu Ser
 20 25 30
 Gly Lys Pro Ala Ile Ile Pro Asp Arg Glu Val Leu Tyr Arg Glu Phe
 35 40 45
 Asp Glu Met Glu Glu Cys
 50

<210> 31
 <211> 686
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Mutant Hepatitis C virus NS3/4A

<400> 31
 Met Ala Pro Ile Thr Ala Tyr Ala Gln Gln Thr Arg Gly Leu Leu Gly
 1 5 10 15
 Cys Ile Ile Thr Ser Leu Thr Gly Arg Asp Lys Asn Gln Val Glu Gly
 20 25 30
 Glu Val Gln Ile Val Ser Thr Ala Ala Gln Thr Phe Leu Ala Thr Cys
 35 40 45
 Ile Asn Gly Val Cys Trp Thr Val Tyr His Gly Ala Gly Thr Arg Thr
 50 55 60
 Ile Ala Ser Pro Lys Gly Pro Val Ile Gln Met Tyr Thr Asn Val Asp
 65 70 75 80
 Gln Asp Leu Val Gly Trp Pro Ala Pro Gln Gly Ala Arg Ser Leu Thr

His	Leu	Glu	Phe	Trp	Glu	Gly	Val	Phe	Thr	Gly	Leu	Thr	His	Ile	Asp
530						535					540				
Ala	His	Phe	Leu	Ser	Gln	Thr	Lys	Gln	Ser	Gly	Glu	Asn	Leu	Pro	Tyr
545					550					555					560
Leu	Val	Ala	Tyr	Gln	Ala	Thr	Val	Cys	Ala	Arg	Ala	Gln	Ala	Pro	Pro
				565						570					575
Pro	Ser	Trp	Asp	Gln	Met	Trp	Lys	Cys	Leu	Ile	Arg	Leu	Lys	Pro	Thr
			580					585					590		
Leu	His	Gly	Pro	Thr	Pro	Leu	Leu	Tyr	Arg	Leu	Gly	Ala	Val	Gln	Asn
		595					600					605			
Glu	Val	Thr	Leu	Thr	His	Pro	Val	Thr	Lys	Tyr	Ile	Met	Thr	Cys	Met
		610				615					620				
Ser	Ala	Asp	Leu	Glu	Val	Val	Thr	Gly	Thr	Trp	Val	Leu	Val	Gly	Gly
625					630					635					640
Val	Leu	Ala	Ala	Leu	Ala	Ala	Tyr	Cys	Leu	Ser	Thr	Gly	Cys	Val	Val
				645					650						655
Ile	Val	Gly	Arg	Ile	Val	Leu	Ser	Gly	Lys	Pro	Ala	Ile	Ile	Pro	Asp
			660					665					670		
Arg	Glu	Val	Leu	Tyr	Arg	Glu	Phe	Asp	Glu	Met	Glu	Glu	Cys		
		675					680					685			

<210> 32

<211> 686

<212> PRT

<213> Artificial Sequence

<220>

<223> Mutant Hepatitis C virus NS3/4A

<400> 32

Met	Ala	Pro	Ile	Thr	Ala	Tyr	Ala	Gln	Gln	Thr	Arg	Gly	Leu	Leu	Gly
1				5					10					15	
Cys	Ile	Ile	Thr	Ser	Leu	Thr	Gly	Arg	Asp	Lys	Asn	Gln	Val	Glu	Gly
			20					25					30		
Glu	Val	Gln	Ile	Val	Ser	Thr	Ala	Ala	Gln	Thr	Phe	Leu	Ala	Thr	Cys
		35					40					45			
Ile	Asn	Gly	Val	Cys	Trp	Thr	Val	Tyr	His	Gly	Ala	Gly	Thr	Arg	Thr
	50					55				60					
Ile	Ala	Ser	Pro	Lys	Gly	Pro	Val	Ile	Gln	Met	Tyr	Thr	Asn	Val	Asp
65				70						75					80
Gln	Asp	Leu	Val	Gly	Trp	Pro	Ala	Pro	Gln	Gly	Ala	Arg	Ser	Leu	Thr
			85						90					95	
Pro	Cys	Thr	Cys	Gly	Ser	Ser	Asp	Leu	Tyr	Leu	Val	Thr	Arg	His	Ala
		100					105						110		
Asp	Val	Ile	Pro	Val	Arg	Arg	Arg	Gly	Asp	Gly	Arg	Gly	Ser	Leu	Leu
		115					120					125			
Ser	Pro	Arg	Pro	Ile	Ser	Tyr	Leu	Lys	Gly	Ser	Ser	Gly	Gly	Pro	Leu
		130				135					140				
Leu	Cys	Pro	Ala	Gly	His	Ala	Val	Gly	Ile	Phe	Arg	Ala	Ala	Val	Cys
145					150					155					160
Thr	Arg	Gly	Val	Ala	Lys	Ala	Val	Asp	Phe	Ile	Pro	Val	Glu	Ser	Leu
			165						170					175	
Glu	Thr	Thr	Met	Arg	Ser	Pro	Val	Phe	Ser	Asp	Asn	Ser	Ser	Pro	Pro
			180					185						190	

Ala Val Pro Gln Ser Tyr Gln Val Ala His Leu His Ala Pro Thr Gly
 195 200 205
 Ser Gly Lys Ser Thr Lys Val Pro Ala Ala Tyr Ala Ala Gln Gly Tyr
 210 215 220
 Lys Val Leu Val Leu Asn Pro Ser Val Ala Ala Thr Met Gly Phe Gly
 225 230 235 240
 Ala Tyr Met Ser Lys Ala His Gly Ile Asp Pro Asn Ile Arg Thr Gly
 245 250 255
 Val Arg Thr Ile Thr Thr Gly Ser Pro Ile Thr Tyr Ser Thr Tyr Gly
 260 265 270
 Lys Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Ile Ile
 275 280 285
 Ile Cys Asp Glu Cys His Ser Thr Asp Ala Thr Ser Ile Leu Gly Ile
 290 295 300
 Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Thr Val
 305 310 315 320
 Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Val Pro His Pro Asn
 325 330 335
 Ile Glu Glu Val Ala Leu Ser Thr Thr Gly Glu Ile Pro Phe Tyr Gly
 340 345 350
 Lys Ala Ile Pro Leu Glu Ala Ile Lys Gly Gly Arg His Leu Ile Phe
 355 360 365
 Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala Ala Lys Leu Val Ala
 370 375 380
 Leu Gly Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ser Val
 385 390 395 400
 Ile Pro Thr Ser Gly Asp Val Val Val Val Ala Thr Asp Ala Leu Met
 405 410 415
 Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Thr Cys
 420 425 430
 Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile Glu
 435 440 445
 Thr Ile Thr Leu Pro Gln Asp Ala Val Ser Arg Thr Gln Arg Arg Gly
 450 455 460
 Arg Thr Gly Arg Gly Lys Pro Gly Ile Tyr Arg Phe Val Ala Pro Gly
 465 470 475 480
 Glu Arg Pro Ser Gly Met Phe Asp Ser Ser Val Leu Cys Glu Cys Tyr
 485 490 495
 Asp Ala Gly Cys Ala Trp Tyr Glu Leu Thr Pro Ala Glu Thr Thr Val
 500 505 510
 Arg Leu Arg Ala Tyr Met Asn Thr Pro Gly Leu Pro Val Cys Gln Asp
 515 520 525
 His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr His Ile Asp
 530 535 540
 Ala His Phe Leu Ser Gln Thr Lys Gln Ser Gly Glu Asn Leu Pro Tyr
 545 550 555 560
 Leu Val Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala Gln Ala Pro Pro
 565 570 575
 Pro Ser Trp Asp Gln Met Trp Lys Cys Leu Ile Arg Leu Lys Pro Thr
 580 585 590
 Leu His Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Ala Val Gln Asn
 595 600 605
 Glu Val Thr Leu Thr His Pro Val Thr Lys Tyr Ile Met Thr Cys Met
 610 615 620
 Ser Ala Asp Leu Glu Val Val Arg Gly Thr Trp Val Leu Val Gly Gly

625		630		635		640									
Val	Leu	Ala	Ala	Leu	Ala	Ala	Tyr	Cys	Leu	Ser	Thr	Gly	Cys	Val	Val
		645		650		655									
Ile	Val	Gly	Arg	Ile	Val	Leu	Ser	Gly	Lys	Pro	Ala	Ile	Ile	Pro	Asp
		660		665		670									
Arg	Glu	Val	Leu	Tyr	Arg	Glu	Phe	Asp	Glu	Met	Glu	Glu	Cys		
		675		680		685									

<210> 33
 <211> 25
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Mutant Hepatitis C virus NS3/4A peptide

<400> 33
Thr Lys Tyr Met Thr Cys Met Ser Ala Asp Leu Glu Val Val Thr Pro
1 5 10 15
Thr Trp Val Leu Val Gly Gly Val Leu
20 25

<210> 34
 <211> 25
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Mutant Hepatitis C virus NS3/4A peptide

<400> 34
Thr Lys Tyr Met Thr Cys Met Ser Ala Asp Leu Glu Val Val Arg Pro
1 5 10 15
Thr Trp Val Leu Val Gly Gly Val Leu
20 25

<210> 35
 <211> 25
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Mutant Hepatitis C virus NS3/4A peptide

<400> 35
Thr Lys Tyr Met Thr Cys Met Ser Ala Asp Leu Glu Val Val Arg Pro
1 5 10 15
Ala Trp Val Leu Val Gly Gly Val Leu
20 25

<210> 36

<211> 25
<212> PRT
<213> Artificial Sequence

<220>
<223> Mutant Hepatitis C virus NS3/4A peptide

<400> 36
Thr Lys Tyr Met Thr Cys Met Ser Ala Asp Leu Glu Val Val Cys Ser
1 5 10 15
Thr Trp Val Leu Val Gly Gly Val Leu
20 25

<210> 37
<211> 25
<212> PRT
<213> Artificial Sequence

<220>
<223> Mutant Hepatitis C virus NS3/4A peptide

<400> 37
Thr Lys Tyr Met Thr Cys Met Ser Ala Asp Leu Glu Val Cys Cys Ser
1 5 10 15
Thr Trp Val Leu Val Gly Gly Val Leu
20 25

<210> 38
<211> 25
<212> PRT
<213> Artificial Sequence

<220>
<223> Mutant Hepatitis C virus NS3/4A peptide

<400> 38
Thr Lys Tyr Met Thr Cys Met Ser Ala Asp Leu Glu Val Ser Ser Ser
1 5 10 15
Thr Trp Val Leu Val Gly Gly Val Leu
20 25

<210> 39
<211> 25
<212> PRT
<213> Artificial Sequence

<220>
<223> Mutant Hepatitis C virus NS3/4A peptide

<400> 39
Thr Lys Tyr Met Thr Cys Met Ser Ala Asp Ser Ser Ser Ser Cys Ser
1 5 10 15

Thr Trp Val Leu Val Gly Gly Val Leu
 20 25

<210> 40
 <211> 25
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Mutant Hepatitis C virus NS3/4A peptide

<400> 40
 Thr Lys Tyr Met Thr Cys Met Ser Ala Asp Val Val Val Val Thr Ser
 1 5 10 15
 Thr Trp Val Leu Val Gly Gly Val Leu
 20 25

<210> 41
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hepatitis C virus NS5 peptide

<400> 41
 Ala Ser Glu Asp Val Val Cys Cys Ser Met Ser Tyr Thr Trp Thr Gly
 1 5 10 15

<210> 42
 <211> 18
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Mutant Hepatitis C virus NS5A/B peptide

<400> 42
 Ser Ser Glu Asp Val Val Cys Cys Ser Met Trp Val Leu Val Gly Gly
 1 5 10 15
 Val Leu

<210> 43
 <211> 686
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Mutant Hepatitis C virus NS3/4A

[illegible][illegible]

Pro	Cys	Thr	Cys	Gly	Ser	Ser	Asp	Leu	Tyr	Leu	Val	Thr	Arg	His	Ala		
			100					105					110				
Asp	Val	Ile	Pro	Val	Arg	Arg	Arg	Gly	Asp	Gly	Arg	Gly	Ser	Leu	Leu		
		115					120					125					
Ser	Pro	Arg	Pro	Ile	Ser	Tyr	Leu	Lys	Gly	Ser	Ser	Gly	Gly	Pro	Leu		
		130				135					140						
Leu	Cys	Pro	Ala	Gly	His	Ala	Val	Gly	Ile	Phe	Arg	Ala	Ala	Val	Cys		
145					150					155					160		
Thr	Arg	Gly	Val	Ala	Lys	Ala	Val	Asp	Phe	Ile	Pro	Val	Glu	Ser	Leu		
				165					170					175			
Glu	Thr	Thr	Met	Arg	Ser	Pro	Val	Phe	Ser	Asp	Asn	Ser	Ser	Pro	Pro		
			180					185					190				
Ala	Val	Pro	Gln	Ser	Tyr	Gln	Val	Ala	His	Leu	His	Ala	Pro	Thr	Gly		
		195					200					205					
Ser	Gly	Lys	Ser	Thr	Lys	Val	Pro	Ala	Ala	Tyr	Ala	Ala	Gln	Gly	Tyr		
		210				215					220						
Lys	Val	Leu	Val	Leu	Asn	Pro	Ser	Val	Ala	Ala	Thr	Met	Gly	Phe	Gly		
225					230					235					240		
Ala	Tyr	Met	Ser	Lys	Ala	His	Gly	Ile	Asp	Pro	Asn	Ile	Arg	Thr	Gly		
				245					250					255			
Val	Arg	Thr	Ile	Thr	Thr	Gly	Ser	Pro	Ile	Thr	Tyr	Ser	Thr	Tyr	Gly		
			260					265					270				
Lys	Phe	Leu	Ala	Asp	Gly	Gly	Cys	Ser	Gly	Gly	Ala	Tyr	Asp	Ile	Ile		
		275					280					285					
Ile	Cys	Asp	Glu	Cys	His	Ser	Thr	Asp	Ala	Thr	Ser	Ile	Leu	Gly	Ile		
		290				295					300						
Gly	Thr	Val	Leu	Asp	Gln	Ala	Glu	Thr	Ala	Gly	Ala	Arg	Leu	Thr	Val		
305					310					315					320		
Leu	Ala	Thr	Ala	Thr	Pro	Pro	Gly	Ser	Val	Thr	Val	Pro	His	Pro	Asn		
				325					330					335			
Ile	Glu	Glu	Val	Ala	Leu	Ser	Thr	Thr	Gly	Glu	Ile	Pro	Phe	Tyr	Gly		
			340					345					350				
Lys	Ala	Ile	Pro	Leu	Glu	Ala	Ile	Lys	Gly	Gly	Arg	His	Leu	Ile	Phe		
		355					360					365					
Cys	His	Ser	Lys	Lys	Lys	Cys	Asp	Glu	Leu	Ala	Ala	Lys	Leu	Val	Ala		
		370				375					380						
Leu	Gly	Val	Asn	Ala	Val	Ala	Tyr	Tyr	Arg	Gly	Leu	Asp	Val	Ser	Val		
385					390					395					400		
Ile	Pro	Thr	Ser	Gly	Asp	Val	Val	Val	Val	Ala	Thr	Asp	Ala	Leu	Met		
				405					410					415			
Thr	Gly	Phe	Thr	Gly	Asp	Phe	Asp	Ser	Val	Ile	Asp	Cys	Asn	Thr	Cys		
			420				425						430				
Val	Thr	Gln	Thr	Val	Asp	Phe	Ser	Leu	Asp	Pro	Thr	Phe	Thr	Ile	Glu		
		435					440					445					
Thr	Ile	Thr	Leu	Pro	Gln	Asp	Ala	Val	Ser	Arg	Thr	Gln	Arg	Arg	Gly		
		450				455					460						
Arg	Thr	Gly	Arg	Gly	Lys	Pro	Gly	Ile	Tyr	Arg	Phe	Val	Ala	Pro	Gly		
465					470					475					480		
Glu	Arg	Pro	Ser	Gly	Met	Phe	Asp	Ser	Ser	Val	Leu	Cys	Glu	Cys	Tyr		
				485					490					495			
Asp	Ala	Gly	Cys	Ala	Trp	Tyr	Glu	Leu	Thr	Pro	Ala	Glu	Thr	Thr	Val		
			500					505					510				
Arg	Leu	Arg	Ala	Tyr	Met	Asn	Thr	Pro	Gly	Leu	Pro	Val	Cys	Gln	Asp		
		515					520					525					
His	Leu	Glu	Phe	Trp	Glu	Gly	Val	Phe	Thr	Gly	Leu	Thr	His	Ile	Asp		

530	535	540
Ala His Phe Leu Ser Gln Thr Lys Gln Ser Gly Glu Asn Leu Pro Tyr		
545	550	555
Leu Val Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala Gln Ala Pro Pro		
	565	570
Pro Ser Trp Asp Gln Met Trp Lys Cys Leu Ile Arg Leu Lys Pro Thr		
	580	585
Leu His Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Ala Val Gln Asn		
	595	600
Glu Val Thr Leu Thr His Pro Val Thr Lys Tyr Ile Met Thr Cys Met		
	610	615
Ser Ala Asp Leu Glu Val Val Arg Pro Thr Trp Val Leu Val Gly Gly		
625	630	635
Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Thr Gly Cys Val Val		
	645	650
Ile Val Gly Arg Ile Val Leu Ser Gly Lys Pro Ala Ile Ile Pro Asp		
	660	665
Arg Glu Val Leu Tyr Arg Glu Phe Asp Glu Met Glu Glu Cys		
	675	680
		685

<210> 45

<211> 686

<212> PRT

<213> Artificial Sequence

<220>

<223> Mutant Hepatitis C virus NS3/4A

<400> 45

Met Ala Pro Ile Thr Ala Tyr Ala Gln Gln Thr Arg Gly Leu Leu Gly		
1	5	10
Cys Ile Ile Thr Ser Leu Thr Gly Arg Asp Lys Asn Gln Val Glu Gly		
	20	25
Glu Val Gln Ile Val Ser Thr Ala Ala Gln Thr Phe Leu Ala Thr Cys		
	35	40
Ile Asn Gly Val Cys Trp Thr Val Tyr His Gly Ala Gly Thr Arg Thr		
	50	55
Ile Ala Ser Pro Lys Gly Pro Val Ile Gln Met Tyr Thr Asn Val Asp		
65	70	75
Gln Asp Leu Val Gly Trp Pro Ala Pro Gln Gly Ala Arg Ser Leu Thr		
	85	90
Pro Cys Thr Cys Gly Ser Ser Asp Leu Tyr Leu Val Thr Arg His Ala		
	100	105
Asp Val Ile Pro Val Arg Arg Arg Gly Asp Gly Arg Gly Ser Leu Leu		
	115	120
Ser Pro Arg Pro Ile Ser Tyr Leu Lys Gly Ser Ser Gly Gly Pro Leu		
	130	135
Leu Cys Pro Ala Gly His Ala Val Gly Ile Phe Arg Ala Ala Val Cys		
145	150	155
Thr Arg Gly Val Ala Lys Ala Val Asp Phe Ile Pro Val Glu Ser Leu		
	165	170
Glu Thr Thr Met Arg Ser Pro Val Phe Ser Asp Asn Ser Ser Pro Pro		
	180	185
Ala Val Pro Gln Ser Tyr Gln Val Ala His Leu His Ala Pro Thr Gly		
		190

Ser Gly Lys	Ser Thr Lys	Val Pro Ala	Ala Tyr Ala	Ala Gln Gly	Tyr
Lys Val Leu	Val Leu Asn	Pro Ser Val	Ala Ala Thr	Met Gly Phe	Gly
Ala Tyr Met	Ser Lys Ala	His Gly Ile	Asp Pro Asn	Ile Arg Thr	Gly
Val Arg Thr	Ile Thr Thr	Gly Ser Pro	Ile Thr Tyr	Ser Thr Tyr	Gly
Lys Phe Leu	Ala Asp Gly	Gly Cys Ser	Gly Gly Ala	Tyr Asp Ile	Ile
Ile Cys Asp	Glu Cys His	Ser Thr Asp	Ala Thr Ser	Ile Leu Gly	Ile
Gly Thr Val	Leu Asp Gln	Ala Glu Thr	Ala Gly Ala	Arg Leu Thr	Val
Leu Ala Thr	Ala Thr Pro	Pro Gly Ser	Val Thr Val	Pro His Pro	Asn
Ile Glu Glu	Val Ala Leu	Ser Thr Thr	Gly Glu Ile	Pro Phe Tyr	Gly
Lys Ala Ile	Pro Leu Glu	Ala Ile Lys	Gly Gly Arg	His Leu Ile	Phe
Cys His Ser	Lys Lys Lys	Cys Asp Glu	Leu Ala Ala	Lys Leu Val	Ala
Leu Gly Val	Asn Ala Val	Ala Tyr Tyr	Arg Gly Leu	Asp Val Ser	Val
Ile Pro Thr	Ser Gly Asp	Val Val Val	Val Ala Thr	Asp Ala Leu	Met
Thr Gly Phe	Thr Gly Asp	Phe Asp Ser	Val Ile Asp	Cys Asn Thr	Cys
Val Thr Gln	Thr Val Asp	Phe Ser Leu	Asp Pro Thr	Phe Thr Ile	Glu
Thr Ile Thr	Leu Pro Gln	Asp Ala Val	Ser Arg Thr	Gln Arg Arg	Gly
Arg Thr Gly	Arg Gly Lys	Pro Gly Ile	Tyr Arg Phe	Val Ala Pro	Gly
Glu Arg Pro	Ser Gly Met	Phe Asp Ser	Ser Val Leu	Cys Glu Cys	Tyr
Asp Ala Gly	Cys Ala Trp	Tyr Glu Leu	Thr Pro Ala	Glu Thr Thr	Val
Arg Leu Arg	Ala Tyr Met	Asn Thr Pro	Gly Leu Pro	Val Cys Gln	Asp
His Leu Glu	Phe Trp Glu	Gly Val Phe	Thr Gly Leu	Thr His Ile	Asp
Ala His Phe	Leu Ser Gln	Thr Lys Gln	Ser Gly Glu	Asn Leu Pro	Tyr
Leu Val Ala	Tyr Gln Ala	Thr Val Cys	Ala Arg Ala	Gln Ala Pro	Pro
Pro Ser Trp	Asp Gln Met	Trp Lys Cys	Leu Ile Arg	Leu Lys Pro	Thr
Leu His Gly	Pro Thr Pro	Leu Leu Tyr	Arg Leu Gly	Ala Val Gln	Asn
Glu Val Thr	Leu Thr His	Pro Val Thr	Lys Tyr Ile	Met Thr Cys	Met
Ser Ala Asp	Leu Glu Val	Val Arg Pro	Ala Trp Val	Leu Val Gly	Gly

Val	Leu	Ala	Ala	Leu	Ala	Ala	Tyr	Cys	Leu	Ser	Thr	Gly	Cys	Val	Val
				645					650					655	
Ile	Val	Gly	Arg	Ile	Val	Leu	Ser	Gly	Lys	Pro	Ala	Ile	Ile	Pro	Asp
			660					665					670		
Arg	Glu	Val	Leu	Tyr	Arg	Glu	Phe	Asp	Glu	Met	Glu	Glu	Cys		
		675					680					685			

<210> 46

<211> 686

<212> PRT

<213> Artificial Sequence

<220>

<223> Mutant Hepatitis C virus NS3/4A

<400> 46

Met	Ala	Pro	Ile	Thr	Ala	Tyr	Ala	Gln	Gln	Thr	Arg	Gly	Leu	Leu	Gly
1				5				10					15		
Cys	Ile	Ile	Thr	Ser	Leu	Thr	Gly	Arg	Asp	Lys	Asn	Gln	Val	Glu	Gly
			20				25					30			
Glu	Val	Gln	Ile	Val	Ser	Thr	Ala	Ala	Gln	Thr	Phe	Leu	Ala	Thr	Cys
		35				40					45				
Ile	Asn	Gly	Val	Cys	Trp	Thr	Val	Tyr	His	Gly	Ala	Gly	Thr	Arg	Thr
	50				55					60					
Ile	Ala	Ser	Pro	Lys	Gly	Pro	Val	Ile	Gln	Met	Tyr	Thr	Asn	Val	Asp
65				70				75						80	
Gln	Asp	Leu	Val	Gly	Trp	Pro	Ala	Pro	Gln	Gly	Ala	Arg	Ser	Leu	Thr
			85			90							95		
Pro	Cys	Thr	Cys	Gly	Ser	Ser	Asp	Leu	Tyr	Leu	Val	Thr	Arg	His	Ala
			100			105						110			
Asp	Val	Ile	Pro	Val	Arg	Arg	Arg	Gly	Asp	Gly	Arg	Gly	Ser	Leu	Leu
	115				120						125				
Ser	Pro	Arg	Pro	Ile	Ser	Tyr	Leu	Lys	Gly	Ser	Ser	Gly	Gly	Pro	Leu
	130				135					140					
Leu	Cys	Pro	Ala	Gly	His	Ala	Val	Gly	Ile	Phe	Arg	Ala	Ala	Val	Cys
145				150					155					160	
Thr	Arg	Gly	Val	Ala	Lys	Ala	Val	Asp	Phe	Ile	Pro	Val	Glu	Ser	Leu
			165			170							175		
Glu	Thr	Thr	Met	Arg	Ser	Pro	Val	Phe	Ser	Asp	Asn	Ser	Ser	Pro	Pro
		180				185						190			
Ala	Val	Pro	Gln	Ser	Tyr	Gln	Val	Ala	His	Leu	His	Ala	Pro	Thr	Gly
	195				200					205					
Ser	Gly	Lys	Ser	Thr	Lys	Val	Pro	Ala	Ala	Tyr	Ala	Ala	Gln	Gly	Tyr
	210				215					220					
Lys	Val	Leu	Val	Leu	Asn	Pro	Ser	Val	Ala	Ala	Thr	Met	Gly	Phe	Gly
225				230					235					240	
Ala	Tyr	Met	Ser	Lys	Ala	His	Gly	Ile	Asp	Pro	Asn	Ile	Arg	Thr	Gly
			245			250							255		
Val	Arg	Thr	Ile	Thr	Thr	Gly	Ser	Pro	Ile	Thr	Tyr	Ser	Thr	Tyr	Gly
		260				265						270			
Lys	Phe	Leu	Ala	Asp	Gly	Gly	Cys	Ser	Gly	Gly	Ala	Tyr	Asp	Ile	Ile
	275				280						285				
Ile	Cys	Asp	Glu	Cys	His	Ser	Thr	Asp	Ala	Thr	Ser	Ile	Leu	Gly	Ile
	290				295					300					

Gly	Thr	Val	Leu	Asp	Gln	Ala	Glu	Thr	Ala	Gly	Ala	Arg	Leu	Thr	Val	305	310	315	320
Leu	Ala	Thr	Ala	Thr	Pro	Pro	Gly	Ser	Val	Thr	Val	Pro	His	Pro	Asn	325	330	335	
Ile	Glu	Glu	Val	Ala	Leu	Ser	Thr	Thr	Gly	Glu	Ile	Pro	Phe	Tyr	Gly	340	345	350	
Lys	Ala	Ile	Pro	Leu	Glu	Ala	Ile	Lys	Gly	Gly	Arg	His	Leu	Ile	Phe	355	360	365	
Cys	His	Ser	Lys	Lys	Lys	Cys	Asp	Glu	Leu	Ala	Ala	Lys	Leu	Val	Ala	370	375	380	
Leu	Gly	Val	Asn	Ala	Val	Ala	Tyr	Tyr	Arg	Gly	Leu	Asp	Val	Ser	Val	385	390	395	400
Ile	Pro	Thr	Ser	Gly	Asp	Val	Val	Val	Val	Ala	Thr	Asp	Ala	Leu	Met	405	410	415	
Thr	Gly	Phe	Thr	Gly	Asp	Phe	Asp	Ser	Val	Ile	Asp	Cys	Asn	Thr	Cys	420	425	430	
Val	Thr	Gln	Thr	Val	Asp	Phe	Ser	Leu	Asp	Pro	Thr	Phe	Thr	Ile	Glu	435	440	445	
Thr	Ile	Thr	Leu	Pro	Gln	Asp	Ala	Val	Ser	Arg	Thr	Gln	Arg	Arg	Gly	450	455	460	
Arg	Thr	Gly	Arg	Gly	Lys	Pro	Gly	Ile	Tyr	Arg	Phe	Val	Ala	Pro	Gly	465	470	475	480
Glu	Arg	Pro	Ser	Gly	Met	Phe	Asp	Ser	Ser	Val	Leu	Cys	Glu	Cys	Tyr	485	490	495	
Asp	Ala	Gly	Cys	Ala	Trp	Tyr	Glu	Leu	Thr	Pro	Ala	Glu	Thr	Thr	Val	500	505	510	
Arg	Leu	Arg	Ala	Tyr	Met	Asn	Thr	Pro	Gly	Leu	Pro	Val	Cys	Gln	Asp	515	520	525	
His	Leu	Glu	Phe	Trp	Glu	Gly	Val	Phe	Thr	Gly	Leu	Thr	His	Ile	Asp	530	535	540	
Ala	His	Phe	Leu	Ser	Gln	Thr	Lys	Gln	Ser	Gly	Glu	Asn	Leu	Pro	Tyr	545	550	555	560
Leu	Val	Ala	Tyr	Gln	Ala	Thr	Val	Cys	Ala	Arg	Ala	Gln	Ala	Pro	Pro	565	570	575	
Pro	Ser	Trp	Asp	Gln	Met	Trp	Lys	Cys	Leu	Ile	Arg	Leu	Lys	Pro	Thr	580	585	590	
Leu	His	Gly	Pro	Thr	Pro	Leu	Leu	Tyr	Arg	Leu	Gly	Ala	Val	Gln	Asn	595	600	605	
Glu	Val	Thr	Leu	Thr	His	Pro	Val	Thr	Lys	Tyr	Ile	Met	Thr	Cys	Met	610	615	620	
Ser	Ala	Asp	Leu	Glu	Val	Val	Cys	Ser	Thr	Trp	Val	Leu	Val	Gly	Gly	625	630	635	640
Val	Leu	Ala	Ala	Leu	Ala	Ala	Tyr	Cys	Leu	Ser	Thr	Gly	Cys	Val	Val	645	650	655	
Ile	Val	Gly	Arg	Ile	Val	Leu	Ser	Gly	Lys	Pro	Ala	Ile	Ile	Pro	Asp	660	665	670	
Arg	Glu	Val	Leu	Tyr	Arg	Glu	Phe	Asp	Glu	Met	Glu	Glu	Cys			675	680	685	

<210> 47

<211> 686

<212> PRT

<213> Artificial Sequence

<220>

<223> Mutant Hepatitis C virus NS3/4A

<400> 47

Met	Ala	Pro	Ile	Thr	Ala	Tyr	Ala	Gln	Gln	Thr	Arg	Gly	Leu	Leu	Gly
1				5				10					15		
Cys	Ile	Ile	Thr	Ser	Leu	Thr	Gly	Arg	Asp	Lys	Asn	Gln	Val	Glu	Gly
			20				25					30			
Glu	Val	Gln	Ile	Val	Ser	Thr	Ala	Ala	Gln	Thr	Phe	Leu	Ala	Thr	Cys
		35					40				45				
Ile	Asn	Gly	Val	Cys	Trp	Thr	Val	Tyr	His	Gly	Ala	Gly	Thr	Arg	Thr
	50					55				60					
Ile	Ala	Ser	Pro	Lys	Gly	Pro	Val	Ile	Gln	Met	Tyr	Thr	Asn	Val	Asp
65				70				75						80	
Gln	Asp	Leu	Val	Gly	Trp	Pro	Ala	Pro	Gln	Gly	Ala	Arg	Ser	Leu	Thr
			85					90					95		
Pro	Cys	Thr	Cys	Gly	Ser	Ser	Asp	Leu	Tyr	Leu	Val	Thr	Arg	His	Ala
			100				105					110			
Asp	Val	Ile	Pro	Val	Arg	Arg	Arg	Gly	Asp	Gly	Arg	Gly	Ser	Leu	Leu
		115				120					125				
Ser	Pro	Arg	Pro	Ile	Ser	Tyr	Leu	Lys	Gly	Ser	Ser	Gly	Gly	Pro	Leu
	130					135				140					
Leu	Cys	Pro	Ala	Gly	His	Ala	Val	Gly	Ile	Phe	Arg	Ala	Ala	Val	Cys
145				150					155					160	
Thr	Arg	Gly	Val	Ala	Lys	Ala	Val	Asp	Phe	Ile	Pro	Val	Glu	Ser	Leu
			165					170					175		
Glu	Thr	Thr	Met	Arg	Ser	Pro	Val	Phe	Ser	Asp	Asn	Ser	Ser	Pro	Pro
			180					185				190			
Ala	Val	Pro	Gln	Ser	Tyr	Gln	Val	Ala	His	Leu	His	Ala	Pro	Thr	Gly
		195				200				205					
Ser	Gly	Lys	Ser	Thr	Lys	Val	Pro	Ala	Ala	Tyr	Ala	Ala	Gln	Gly	Tyr
	210					215				220					
Lys	Val	Leu	Val	Leu	Asn	Pro	Ser	Val	Ala	Ala	Thr	Met	Gly	Phe	Gly
225				230					235					240	
Ala	Tyr	Met	Ser	Lys	Ala	His	Gly	Ile	Asp	Pro	Asn	Ile	Arg	Thr	Gly
			245					250					255		
Val	Arg	Thr	Ile	Thr	Thr	Gly	Ser	Pro	Ile	Thr	Tyr	Ser	Thr	Tyr	Gly
			260					265				270			
Lys	Phe	Leu	Ala	Asp	Gly	Gly	Cys	Ser	Gly	Gly	Ala	Tyr	Asp	Ile	Ile
	275					280					285				
Ile	Cys	Asp	Glu	Cys	His	Ser	Thr	Asp	Ala	Thr	Ser	Ile	Leu	Gly	Ile
	290					295				300					
Gly	Thr	Val	Leu	Asp	Gln	Ala	Glu	Thr	Ala	Gly	Ala	Arg	Leu	Thr	Val
305				310						315				320	
Leu	Ala	Thr	Ala	Thr	Pro	Pro	Gly	Ser	Val	Thr	Val	Pro	His	Pro	Asn
			325					330				335			
Ile	Glu	Glu	Val	Ala	Leu	Ser	Thr	Thr	Gly	Glu	Ile	Pro	Phe	Tyr	Gly
			340					345				350			
Lys	Ala	Ile	Pro	Leu	Glu	Ala	Ile	Lys	Gly	Gly	Arg	His	Leu	Ile	Phe
	355					360					365				
Cys	His	Ser	Lys	Lys	Lys	Cys	Asp	Glu	Leu	Ala	Ala	Lys	Leu	Val	Ala
	370					375					380				
Leu	Gly	Val	Asn	Ala	Val	Ala	Tyr	Tyr	Arg	Gly	Leu	Asp	Val	Ser	Val
385				390						395				400	
Ile	Pro	Thr	Ser	Gly	Asp	Val	Val	Val	Val	Ala	Thr	Asp	Ala	Leu	Met

				405					410				415				
Thr	Gly	Phe	Thr	Gly	Asp	Phe	Asp	Ser	Val	Ile	Asp	Cys	Asn	Thr	Cys		
			420					425					430				
Val	Thr	Gln	Thr	Val	Asp	Phe	Ser	Leu	Asp	Pro	Thr	Phe	Thr	Ile	Glu		
		435					440					445					
Thr	Ile	Thr	Leu	Pro	Gln	Asp	Ala	Val	Ser	Arg	Thr	Gln	Arg	Arg	Gly		
	450					455					460						
Arg	Thr	Gly	Arg	Gly	Lys	Pro	Gly	Ile	Tyr	Arg	Phe	Val	Ala	Pro	Gly		
465					470					475					480		
Glu	Arg	Pro	Ser	Gly	Met	Phe	Asp	Ser	Ser	Val	Leu	Cys	Glu	Cys	Tyr		
			485					490					495				
Asp	Ala	Gly	Cys	Ala	Trp	Tyr	Glu	Leu	Thr	Pro	Ala	Glu	Thr	Thr	Val		
		500						505					510				
Arg	Leu	Arg	Ala	Tyr	Met	Asn	Thr	Pro	Gly	Leu	Pro	Val	Cys	Gln	Asp		
	515					520						525					
His	Leu	Glu	Phe	Trp	Glu	Gly	Val	Phe	Thr	Gly	Leu	Thr	His	Ile	Asp		
	530					535					540						
Ala	His	Phe	Leu	Ser	Gln	Thr	Lys	Gln	Ser	Gly	Glu	Asn	Leu	Pro	Tyr		
545					550					555					560		
Leu	Val	Ala	Tyr	Gln	Ala	Thr	Val	Cys	Ala	Arg	Ala	Gln	Ala	Pro	Pro		
			565					570					575				
Pro	Ser	Trp	Asp	Gln	Met	Trp	Lys	Cys	Leu	Ile	Arg	Leu	Lys	Pro	Thr		
			580					585					590				
Leu	His	Gly	Pro	Thr	Pro	Leu	Leu	Tyr	Arg	Leu	Gly	Ala	Val	Gln	Asn		
	595						600					605					
Glu	Val	Thr	Leu	Thr	His	Pro	Val	Thr	Lys	Tyr	Ile	Met	Thr	Cys	Met		
	610					615					620						
Ser	Ala	Asp	Leu	Glu	Val	Cys	Cys	Ser	Thr	Trp	Val	Leu	Val	Gly	Gly		
625					630					635					640		
Val	Leu	Ala	Ala	Leu	Ala	Ala	Tyr	Cys	Leu	Ser	Thr	Gly	Cys	Val	Val		
			645					650					655				
Ile	Val	Gly	Arg	Ile	Val	Leu	Ser	Gly	Lys	Pro	Ala	Ile	Ile	Pro	Asp		
		660						665					670				
Arg	Glu	Val	Leu	Tyr	Arg	Glu	Phe	Asp	Glu	Met	Glu	Glu	Cys				
	675						680					685					

<210> 48

<211> 686

<212> PRT

<213> Artificial Sequence

<220>

<223> Mutant Hepatitis C virus NS3/4A

<400> 48

Met	Ala	Pro	Ile	Thr	Ala	Tyr	Ala	Gln	Gln	Thr	Arg	Gly	Leu	Leu	Gly		
1				5				10					15				
Cys	Ile	Ile	Thr	Ser	Leu	Thr	Gly	Arg	Asp	Lys	Asn	Gln	Val	Glu	Gly		
			20					25				30					
Glu	Val	Gln	Ile	Val	Ser	Thr	Ala	Ala	Gln	Thr	Phe	Leu	Ala	Thr	Cys		
		35					40				45						
Ile	Asn	Gly	Val	Cys	Trp	Thr	Val	Tyr	His	Gly	Ala	Gly	Thr	Arg	Thr		
	50					55				60							
Ile	Ala	Ser	Pro	Lys	Gly	Pro	Val	Ile	Gln	Met	Tyr	Thr	Asn	Val	Asp		

Arg	Leu	Arg	Ala	Tyr	Met	Asn	Thr	Pro	Gly	Leu	Pro	Val	Cys	Gln	Asp
	515						520					525			
His	Leu	Glu	Phe	Trp	Glu	Gly	Val	Phe	Thr	Gly	Leu	Thr	His	Ile	Asp
	530					535					540				
Ala	His	Phe	Leu	Ser	Gln	Thr	Lys	Gln	Ser	Gly	Glu	Asn	Leu	Pro	Tyr
545					550					555					560
Leu	Val	Ala	Tyr	Gln	Ala	Thr	Val	Cys	Ala	Arg	Ala	Gln	Ala	Pro	Pro
				565					570					575	
Pro	Ser	Trp	Asp	Gln	Met	Trp	Lys	Cys	Leu	Ile	Arg	Leu	Lys	Pro	Thr
			580					585					590		
Leu	His	Gly	Pro	Thr	Pro	Leu	Leu	Tyr	Arg	Leu	Gly	Ala	Val	Gln	Asn
		595					600					605			
Glu	Val	Thr	Leu	Thr	His	Pro	Val	Thr	Lys	Tyr	Ile	Met	Thr	Cys	Met
	610						615					620			
Ser	Ala	Asp	Leu	Glu	Val	Ser	Ser	Ser	Thr	Trp	Val	Leu	Val	Gly	Gly
625					630					635					640
Val	Leu	Ala	Ala	Leu	Ala	Ala	Tyr	Cys	Leu	Ser	Thr	Gly	Cys	Val	Val
				645					650					655	
Ile	Val	Gly	Arg	Ile	Val	Leu	Ser	Gly	Lys	Pro	Ala	Ile	Ile	Pro	Asp
			660					665					670		
Arg	Glu	Val	Leu	Tyr	Arg	Glu	Phe	Asp	Glu	Met	Glu	Glu	Cys		
		675					680					685			

<210> 49

<211> 686

<212> PRT

<213> Artificial Sequence

<220>

<223> Mutant Hepatitis C virus NS3/4A

<400> 49

Met	Ala	Pro	Ile	Thr	Ala	Tyr	Ala	Gln	Gln	Thr	Arg	Gly	Leu	Leu	Gly
1				5				10					15		
Cys	Ile	Ile	Thr	Ser	Leu	Thr	Gly	Arg	Asp	Lys	Asn	Gln	Val	Glu	Gly
			20					25				30			
Glu	Val	Gln	Ile	Val	Ser	Thr	Ala	Ala	Gln	Thr	Phe	Leu	Ala	Thr	Cys
		35					40				45				
Ile	Asn	Gly	Val	Cys	Trp	Thr	Val	Tyr	His	Gly	Ala	Gly	Thr	Arg	Thr
	50					55				60					
Ile	Ala	Ser	Pro	Lys	Gly	Pro	Val	Ile	Gln	Met	Tyr	Thr	Asn	Val	Asp
65				70					75					80	
Gln	Asp	Leu	Val	Gly	Trp	Pro	Ala	Pro	Gln	Gly	Ala	Arg	Ser	Leu	Thr
			85					90					95		
Pro	Cys	Thr	Cys	Gly	Ser	Ser	Asp	Leu	Tyr	Leu	Val	Thr	Arg	His	Ala
		100					105					110			
Asp	Val	Ile	Pro	Val	Arg	Arg	Arg	Gly	Asp	Gly	Arg	Gly	Ser	Leu	Leu
	115						120					125			
Ser	Pro	Arg	Pro	Ile	Ser	Tyr	Leu	Lys	Gly	Ser	Ser	Gly	Gly	Pro	Leu
	130					135					140				
Leu	Cys	Pro	Ala	Gly	His	Ala	Val	Gly	Ile	Phe	Arg	Ala	Ala	Val	Cys
145					150				155						160
Thr	Arg	Gly	Val	Ala	Lys	Ala	Val	Asp	Phe	Ile	Pro	Val	Glu	Ser	Leu
			165					170					175		

610		615		620											
Ser	Ala	Asp	Ser	Ser	Ser	Ser	Cys	Ser	Thr	Trp	Val	Leu	Val	Gly	Gly
625					630					635					640
Val	Leu	Ala	Ala	Leu	Ala	Ala	Tyr	Cys	Leu	Ser	Thr	Gly	Cys	Val	Val
				645						650					655
Ile	Val	Gly	Arg	Ile	Val	Leu	Ser	Gly	Lys	Pro	Ala	Ile	Ile	Pro	Asp
			660					665						670	
Arg	Glu	Val	Leu	Tyr	Arg	Glu	Phe	Asp	Glu	Met	Glu	Glu	Cys		
		675					680						685		